

PATENT
10/023,969
Docket 084/002

AMENDMENT TO SPECIFICATION

[Paragraph beginning on page 13, line 3]

a' SEQ. ID NO:1 is a listing of the sequence data obtained. Nucleotides 1-43 and 15376-15418 are plasmid sequence. Thus, the genomic insert begins at residue 44 and ends at residue 15375. The beginning of the cloned cDNA fragment corresponds to residue 13490. There are Alu sequence elements located -1700 base pairs upstream. The sequence of the hTERT insert of pGRN142 can now be obtained from GenBank (~~http://www.ncbi.nlm.nih.gov/~~ the GenBank website) under Accession PGRN142.INS AF121948. Numbering of hTERT residues for plasmids in the following description begins from the translation initiation codon, according to standard practice in the field. The hTERT ATG codon (the translation initiation site) begins at residue 13545 of SEQ. ID NO:1. Thus, position -1, the first upstream residue, corresponds to nucleotide 13544 in SEQ. ID NO:1.